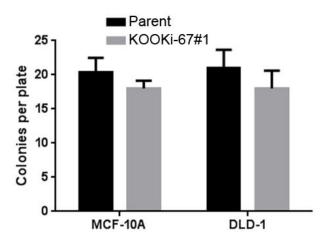
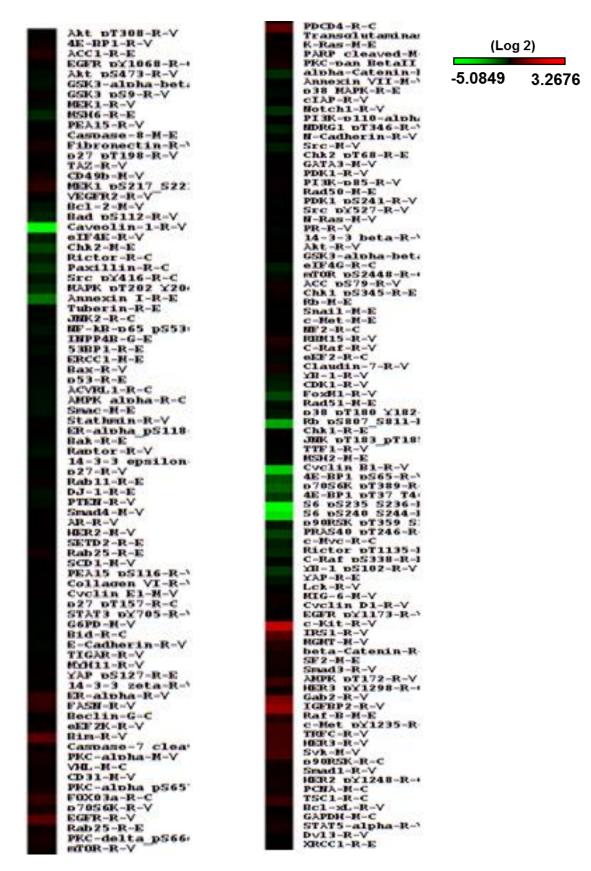
Ki-67 is required for maintenance of cancer stem cells but not cell proliferation

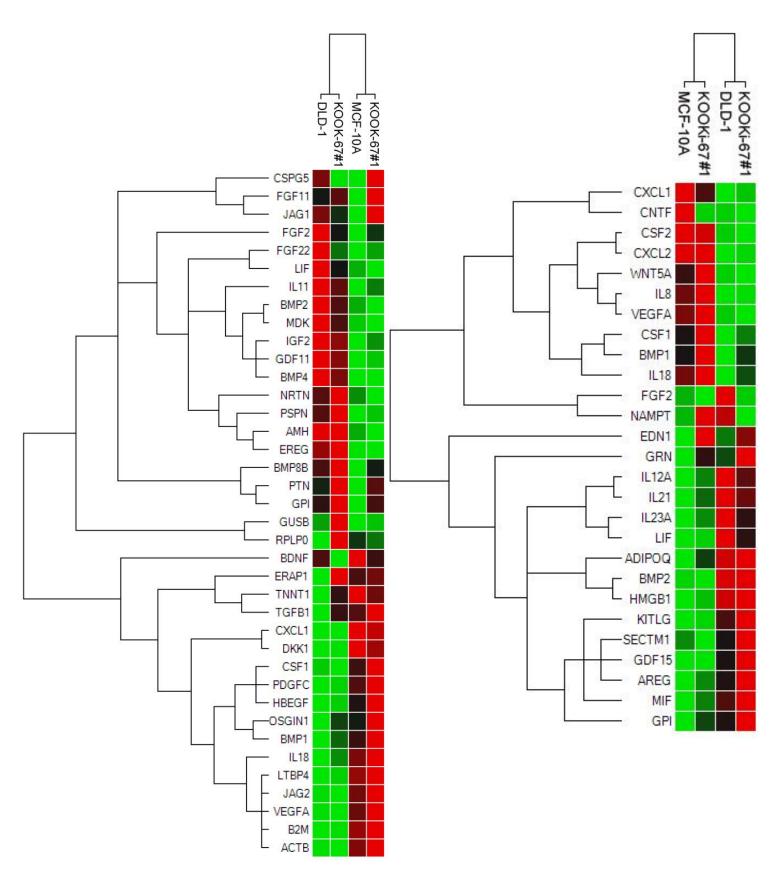
Supplemental Materials



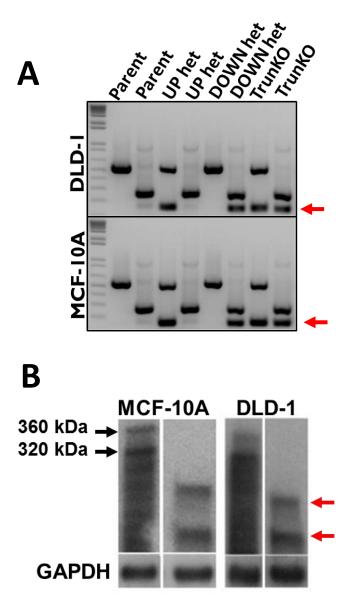
Supplemental Figure 1. Knock out of Ki-67 does not affect colony number. Parental MCF-10A, DLD-1 and derivative KOOKi-67 clones were seeded at 0.5 cell per well in 96-well plates and grown for 30 days under standard conditions, at which time the number of colonies per well were counted.



Supplemental Figure 2: **List of proteins expression analyzed by RPPA.** MCF-10A and DLD-1 parental and KOOKi-67 clones were subjected to RPPA analysis. DLD-1 cells were grown in standard full growth conditions while MCF-10A cells were grown in both full growth (+EGF) and growth arrest (-EGF) conditions.



Supplemental Figure 3. Gene expression of growth factors and cytokines are similar between parental and KOOKi-67 cells. Quantitative PCR was performed using cDNA from parental and KOOKi-67 cells to assay for differences in growth factor and cytokine mRNA expression.



Supplemental Figure 4. PCR and western blot confirmation of TrunKO clones. A) PCR was performed using genomic DNA from parental and TrunKO cells to confirm correct targeting at genomic loci. Each genomic locus was assayed independently using primers specific for the targeted region. The presence of small molecular weight band (red arrow) confirmed targeting at the given locus. **B)** Western blot was performed using lysates from parental MCF-10A and DLD-1 cells (left lanes) and lysates from TrunKO clones (right lanes). Shown are the full length parental isoforms (360 kDa and 320 kDA), as well as the presence of truncated proteins (red arrows). GAPDH was used as a loading control.

Supplemental Table 1: Tumor engraftment is reduced by loss of Ki-67.

Tumor Engraftment							
		Number of cells per injection					
		10^{6}	10 ⁵	10^4	10^{3}		
	Parent	10/10	10/10	9/10	8/10		
Cell Line	KOOKi-67 #1	10/10	9/10	8/10	6/10		
	KOOKi-67 #2	10/10	9/10	8/10	6/10		

Supplemental Table 2: Tumor engraftment is unaffected in Ki-67 TrunKO cells.

Tumor Engraftment							
		Number of cells per injection					
		10^{6}	10 ⁵	104	10^{3}		
Cell Line	Parent	20/20	20/20	19/20	18/19		
	KOOKi-67 #1	10/10	9/10	8/10	7/10		
	KOOKi-67 #2	10/10	9/10	4/5	4/5		
	TrunKO	10/10	9/9	10/10	9/10		

Supplemental Table 3. Cloning and screening primers used in this study.

Cloning Primers							
Region	Forward	Reverse	Screening				
5'HA	GATGCACCCAGGTATTTTCC	CCGCTCCTTTTGATAGTAACC	GTGCCCAGATGTTTGGTCC				
UP 3'HA			CTGAAGCTGTGGGGTTGG				
DOWN 3'HA			CTCATGCAGTTTCAAAACAC				
Region	Forward	Reverse					
UP Allele	GGTTACTATCAAAAGGAGCGG	CACCTGATGCCTGAATGATGG					
DOWN Allele	GGTTACTATCAAAAGGAGCGG	GATGCTAGCACTTCTGTGC					
Deleted Region	eleted Region GTGTCCCAATGTTTGGTCC GGAAAGTGGGGACCGTCGAC						
Cloning Primers							
Region	Forward	Reverse	Screening				
UP 5'HA	CCAGAACCAATAAACACCCC	TAGTTTTGCCAGCAGCCAC	CTCATGAAAGACACGGCACG				
UP 3'HA	AGCAGTAAGTGGTGAGAAAAAC	TAGCAGGTCCAGTTTCTCCAC	CGTCTGTGTGAGCTTGCCG				
DOWN 5'HA	CCAGTCAAAACCCCAACAAGC	TGCTGGTTTGGGTGTGTC	GAAGTATCCTGCAAATCTCC				
DOWN 3'HA GACACCCCAGCAAGCACAAAG GTGTGTGTGTGTTTT		GTGTGTGTGTGCTTTGCC	GCTGGGTTCCTCTTCTACTG				
DOWN 5'HA CCAGTCAAAACCCCAACAAGC TGCTGGTTTGGGTGTCTC GAAGTATCCTGCAAA DOWN 3'HA GACACCCCAGCAAGCACAAAG GTGTGTGTGTGTGTTGCC GCTGGGTTCCTCTTC Knockout Confirmation Screens							
Region	Forward	Reverser					
UP	GTGGCTGCTGGCAAAACTA	GTTTTTCTCACCACTTACTGCT					
DOWN	GACACACCCAAACCAGCA	CTTTGTGCTTGCTGGGGTGTC					
	5'HA UP 3'HA DOWN 3'HA Region UP Allele DOWN Allele Deleted Region UP 5'HA UP 3'HA DOWN 5'HA DOWN 3'HA Region UP	Region Forward 5'HA GATGCACCCAGGTATTTTCC UP 3'HA CACTTTCCCCTGAGCCTCAG DOWN 3'HA GCACAGAAGTGCTAGCATC Knocke Region Forward UP Allele GGTTACTATCAAAAGGAGCGG DOWN Allele GGTTACTATCAAAAGGAGCGG Deleted Region GTGTCCCAATGTTTGGTCC Region Forward UP 5'HA CCAGAACCAATAAACACCCC UP 3'HA AGCAGTAAGTGGTGAGAAAAAC DOWN 5'HA CCAGTCAAAACCCCAACAAGC DOWN 3'HA GACACCCCAGCAACAAGC Region Forward Knocke Region Forward UP GTGGCTGCTGGCAAAACTA	Region Forward Reverse 5'HA GATGCACCCAGGTATTTTCC CCGCTCCTTTTGATAGTAACC UP 3'HA CACTTTCCCCTGAGCCTCAG CACCTGATGCCTGAATGATGG DOWN 3'HA GCACAGAAGTGCTAGCATC GCACCCAAATCCTTGAATTCC Knockout Confirmation Screens Region Forward Reverse UP Allele GGTTACTATCAAAAAGGAGCGG CACCTGATGCCTGAATGATGG DOWN Allele GGTTACTATCAAAAAGGAGCGG GATGCTAGCACTTCTGTGC Deleted Region GTGTCCCAATGTTTGGTCC GGAAAGTGGGGACCGTCGAC Cloning Primers Region Forward Reverse UP 5'HA CCAGAACCAATAAACACCCC TAGTTTTGCCAGCAGCCAC UP 3'HA AGCAGTAAGTGGTGAGAAAAAC TAGCAGGTCCAGTTTCTCCAC DOWN 5'HA CCAGTCAAAACCCCAACAAGC TGCTGGTTTGGTGCC CMN 3'HA GACACCCCAGCAAGAACACAAAG GTGTGTGTGTGTTTGCC Knockout Confirmation Screens Region Forward Reverser UP GTGGCTGCTGGCAAAACTA GTTTTCTCACCACTTACTGCT				